

50704/KMO/B583

SEQUENCE LISTING

(1) INFORMATION FOR SEQ ID NO. 1:

- (i) LENGTH: 365 amino acids
- (ii) MOLECULE TYPE: protein
- (vi) ORIGINAL SOURCE:
 - (a)ORGANISM: Candida boidinii

Met Gly Lys Ile Val Leu Val Leu Tyr Asp Ala Gly Lys His Ala Ala
1 5 10 15

Asp Glu Glu Lys Leu Tyr Gly Cys Thr Glu Asn Lys Leu Gly Ile Ala
20 25 30

Asn Trp Leu Lys Asp Gln Gly His Glu Leu Ile Thr Thr Ser Asp Lys
35 40 45

Glu Gly Glu Thr Ser Glu Leu Asp Lys His Ile Pro Asp Ala Asp Ile
50 55 60

Ile Ile Thr Thr Pro Phe His Pro Ala Tyr Ile Thr Lys Glu Arg Leu
65 70 75 80

Asp Lys Ala Lys Asn Leu Lys Leu Val Val Val Ala Gly Val Gly Ser
85 90 95

Asp His Ile Asp Leu Asp Tyr Ile Asn Gln Thr Gly Lys Lys Ile Ser
100 105 110

Val Leu Glu Val Thr Gly Ser Asn Val Val Ser Val Ala Glu His Val
115 120 125

Val Met Thr Met Leu Val Leu Val Arg Asn Phe Val Pro Ala His Glu
130 135 140

Gln Ile Ile Asn His Asp Trp Glu Val Ala Ala Ile Ala Lys Asp Ala
145 150 155 160

Tyr Asp Ile Glu Gly Lys Thr Ile Ala Thr Ile Gly Ala Gly Arg Ile
165 170 175

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Gly Tyr Arg Val Leu Glu Arg Leu Leu Pro Phe Asn Pro Lys Glu Leu
180 185 190

Leu Tyr Tyr Asp Tyr Gln Ala Leu Pro Lys Glu Ala Glu Glu Lys Val
195 200 205

Gly Ala Arg Arg Val Glu Asn Ile Glu Glu Leu Val Ala Gln Ala Asp
210 215 220

Ile Val Thr Val Asn Ala Pro Leu His Ala Gly Thr Lys Gly Leu Ile
225 230 235 240

Asn Lys Glu Leu Leu Ser Lys Phe Lys Lys Gly Ala Trp Leu Val Asn
245 250 255

Thr Ala Arg Gly Ala Ile Cys Val Ala Glu Asp Val Ala Ala Ala Leu
260 265 270

Glu Ser Gly Gln Leu Arg Gly Tyr Gly Gly Asp Val Trp Phe Pro Gln
275 280 285

Pro Ala Pro Lys Asp His Pro Trp Arg Asp Met Arg Asn Lys Tyr Gly
290 295 300

Ala Gly Asn Ala Met Thr Pro His Tyr Ser Gly Thr Thr Leu Asp Ala
305 310 315 320

Gln Thr Arg Tyr Ala Glu Gly Thr Lys Asn Ile Leu Glu Ser Phe Phe
325 330 335

Thr Gly Lys Phe Asp Tyr Arg Pro Gln Asp Ile Ile Leu Leu Asn Gly
340 345 350

Glu Tyr Val Thr Lys Ala Tyr Gly Lys His Asp Lys Lys
355 360 365

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(2) INFORMATION FOR SEQ ID NO. 2:

- (i) LENGTH: 367 amino acids
- (ii) MOLECULE TYPE: protein
- (vi) ORIGINAL SOURCE:
 - (a) ORGANISM: Candida boidinii

Met Gly Lys Ile Phe Asp Tyr Met Glu Lys Tyr Asp Tyr Glu Gln Leu
1 5 10 15

Val Met Cys Gln Asp Lys Glu Ser Gly Leu Lys Ala Ile Ile Cys Ile
20 25 30

His Val Thr Thr Leu Gly Pro Ala Leu Gly Gly Met Arg Met Trp Thr
35 40 45

Tyr Ala Ser Glu Glu Ala Ile Glu Asp Ala Leu Arg Leu Gly Arg
50 55 60

Gly Met Thr Tyr Lys Asn Ala Ala Gly Leu Asn Leu Gly Gly Gly
65 70 75 80

Lys Thr Val Ile Ile Gly Asp Pro Arg Lys Asp Lys Asn Glu Ala Met
85 90 95

Phe Arg Ala Leu Gly Arg Phe Ile Gln Gly Leu Asn Gly Arg Tyr Ile
100 105 110

Thr Ala Glu Asp Val Gly Thr Thr Val Glu Asp Met Asp Ile Ile His
115 120 125

Glu Glu Thr Arg Tyr Val Thr Gly Val Ser Pro Ala Phe Gly Ser Ser
130 135 140

Gly Asn Pro Ser Pro Val Thr Ala Tyr Gly Val Tyr Arg Gly Met Lys
145 150 155 160

Ala Ala Ala Lys Glu Ala Phe Gly Asp Asp Ser Leu Glu Gly Lys Val
165 170 175

Val Ala Val Gln Gly Val Gly His Val Ala Tyr Glu Leu Cys Lys His
180 185 190

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Leu	His	Asn	Glu	Gly	Ala	Lys	Leu	Ile	Val	Thr	Asp	Ile	Asn	Lys	Glu
195							200					205			
Asn	Ala	Asp	Arg	Ala	Val	Gln	Glu	Phe	Gly	Ala	Glu	Phe	Val	His	Pro
210					215						220				
Asp	Lys	Ile	Tyr	Asp	Val	Glu	Cys	Asp	Ile	Phe	Ala	Pro	Cys	Ala	Leu
225					230				235			240			
Gly	Ala	Ile	Ile	Asn	Asp	Glu	Thr	Ile	Glu	Arg	Leu	Lys	Cys	Lys	Val
245								250				255			
Val	Ala	Gly	Ser	Ala	Asn	Asn	Gln	Leu	Lys	Glu	Glu	Arg	His	Gly	Lys
260								265				270			
Met	Leu	Glu	Glu	Lys	Gly	Ile	Val	Tyr	Ala	Pro	Asp	Tyr	Val	Ile	Asn
275						280					285				
Ala	Gly	Gly	Val	Ile	Asn	Val	Ala	Asp	Glu	Leu	Leu	Gly	Tyr	Asn	Arg
290						295				300					
Glu	Arg	Ala	Met	Lys	Lys	Val	Glu	Gly	Ile	Tyr	Asp	Lys	Ile	Leu	Lys
305						310				315			320		
Val	Phe	Glu	Ile	Ala	Lys	Arg	Asp	Gly	Ile	Pro	Ser	Tyr	Leu	Ala	Ala
325									330			335			
Asp	Arg	Met	Ala	Glu	Glu	Arg	Ile	Glu	Met	Met	Arg	Lys	Thr	Arg	Ser
340								345				350			
Thr	Phe	Leu	Gln	Asp	Gln	Arg	Asn	Leu	Ile	Asn	Phe	Asn	Asn	Lys	
355								360				365			

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(3) INFORMATION FOR SEQ ID NO. 3:

- (i) LENGTH: 640 amino acids
- (ii) MOLECULE TYPE: protein
- (vi) ORIGINAL SOURCE:
 - (a) ORGANISM: *Candida boidinii*

Met Ala Ser Ala Pro Ile Gly Ser Ala Ile Ser Arg Asn Asn Trp Ala
1 5 10 15

Val Thr Cys Asp Ser Ala Gln Ser Gly Asn Glu Cys Asn Lys Ala Ile
20 25 30

Asp Gly Asn Lys Asp Thr Phe Trp His Thr Phe Tyr Gly Ala Asn Gly
35 40 45

Asp Pro Lys Pro Pro His Thr Tyr Thr Ile Asp Met Lys Thr Thr Gln
50 55 60

Asn Val Asn Gly Leu Ser Met Leu Pro Arg Gln Asp Gly Asn Gln Asn
65 70 75 80

Gly Trp Ile Gly Arg His Glu Val Tyr Leu Ser Ser Asp Gly Thr Asn
85 90 95

Trp Gly Ser Pro Val Ala Ser Gly Ser Trp Phe Ala Asp Ser Thr Thr
100 105 110

Lys Tyr Ser Asn Phe Glu Thr Arg Pro Ala Arg Tyr Val Arg Leu Val
115 120 125

Ala Ile Thr Glu Ala Asn Gly Gln Pro Trp Thr Ser Ile Ala Glu Ile
130 135 140

Asn Val Phe Gln Ala Ser Ser Tyr Thr Ala Pro Gln Pro Gly Leu Gly
145 150 155 160

Arg Trp Gly Pro Thr Ile Asp Leu Pro Ile Val Pro Ala Ala Ala Ala
165 170 175

Ile Glu Pro Thr Ser Gly Arg Val Leu Met Trp Ser Ser Tyr Arg Asn
180 185 190

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Asp Ala Phe Gly Gly Ser Pro Gly Gly Ile Thr Leu Thr Ser Ser Trp
195 200 205

Asp Pro Ser Thr Gly Ile Val Ser Asp Arg Thr Val Thr Val Thr Lys
210 215 220

His Asp Met Phe Cys Pro Gly Ile Ser Met Asp Gly Asn Gly Gln Ile
225 230 235 240

Val Val Thr Gly Gly Asn Asp Ala Lys Lys Thr Ser Leu Tyr Asp Ser
245 250 255

Ser Ser Asp Ser Trp Ile Pro Gly Pro Asp Met Gln Val Ala Arg Gly
260 265 270

Tyr Gln Ser Ser Ala Thr Met Ser Asp Gly Arg Val Phe Thr Ile Gly
275 280 285

Gly Ser Trp Ser Gly Gly Val Phe Glu Lys Asn Gly Glu Val Tyr Ser
290 295 300

Pro Ser Ser Lys Thr Trp Thr Ser Leu Pro Asn Ala Lys Val Asn Pro
305 310 315 320

Met Leu Thr Ala Asp Lys Gln Gly Leu Tyr Arg Ser Asp Asn His Ala
325 330 335

Trp Leu Phe Gly Trp Lys Lys Gly Ser Val Phe Gln Ala Gly Pro Ser
340 345 350

Thr Ala Met Asn Trp Tyr Tyr Thr Ser Gly Ser Gly Asp Val Lys Ser
355 360 365

Ala Gly Lys Arg Gln Ser Asn Arg Gly Val Ala Pro Asp Ala Met Cys
370 375 380

Gly Asn Ala Val Met Tyr Asp Ala Val Lys Gly Lys Ile Leu Thr Phe
385 390 395 400

Gly Gly Ser Pro Asp Tyr Gln Asp Ser Asp Ala Thr Thr Asn Ala His
405 410 415

Ile Ile Thr Leu Gly Glu Pro Gly Thr Ser Pro Asn Thr Val Phe Ala
420 425 430

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Ser Asn Gly Leu Tyr Phe Ala Arg Thr Phe His Thr Ser Val Val Leu
435 440 445

Pro Asp Gly Ser Thr Phe Ile Thr Gly Gly Gln Arg Arg Gly Ile Pro
450 455 460

Phe Glu Asp Ser Thr Pro Val Phe Thr Pro Glu Ile Tyr Val Pro Glu
465 470 475 480

Gln Asp Thr Phe Tyr Lys Gln Asn Pro Asn Ser Ile Val Arg Val Tyr
485 490 495

His Ser Ile Ser Leu Leu Leu Pro Asp Gly Arg Val Phe Asn Gly Gly
500 505 510

Gly Gly Leu Cys Gly Asp Cys Thr Thr Asn His Phe Asp Ala Gln Ile
515 520 525

Phe Thr Pro Asn Tyr Leu Tyr Asn Ser Asn Gly Asn Leu Ala Thr Arg
530 535 540

Pro Lys Ile Thr Arg Thr Ser Thr Gln Ser Val Lys Val Gly Gly Arg
545 550 555 560

Ile Thr Ile Ser Thr Asp Ser Ser Ile Ser Lys Ala Ser Leu Ile Arg
565 570 575

Tyr Gly Thr Ala Thr His Thr Val Asn Thr Asp Gln Arg Arg Ile Pro
580 585 590

Leu Thr Leu Thr Asn Asn Gly Gly Asn Ser Tyr Ser Phe Gln Val Pro
595 600 605

Ser Asp Ser Gly Val Ala Leu Pro Gly Tyr Trp Met Leu Phe Val Met
610 615 620

Asn Ser Ala Gly Val Pro Ser Val Ala Ser Thr Ile Arg Val Thr Gln
625 630 635 640